

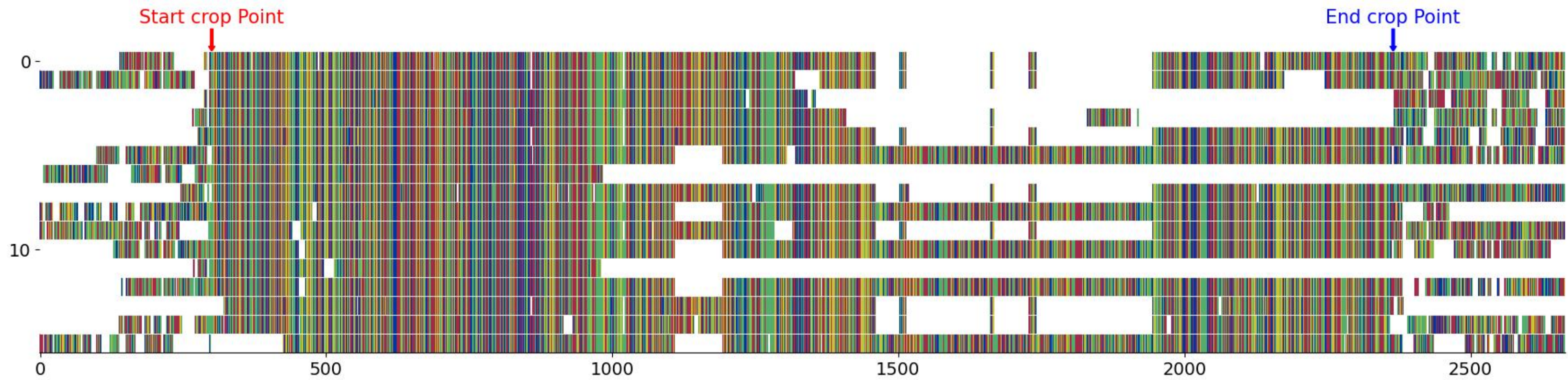
End crop Point

MSA length = 2066



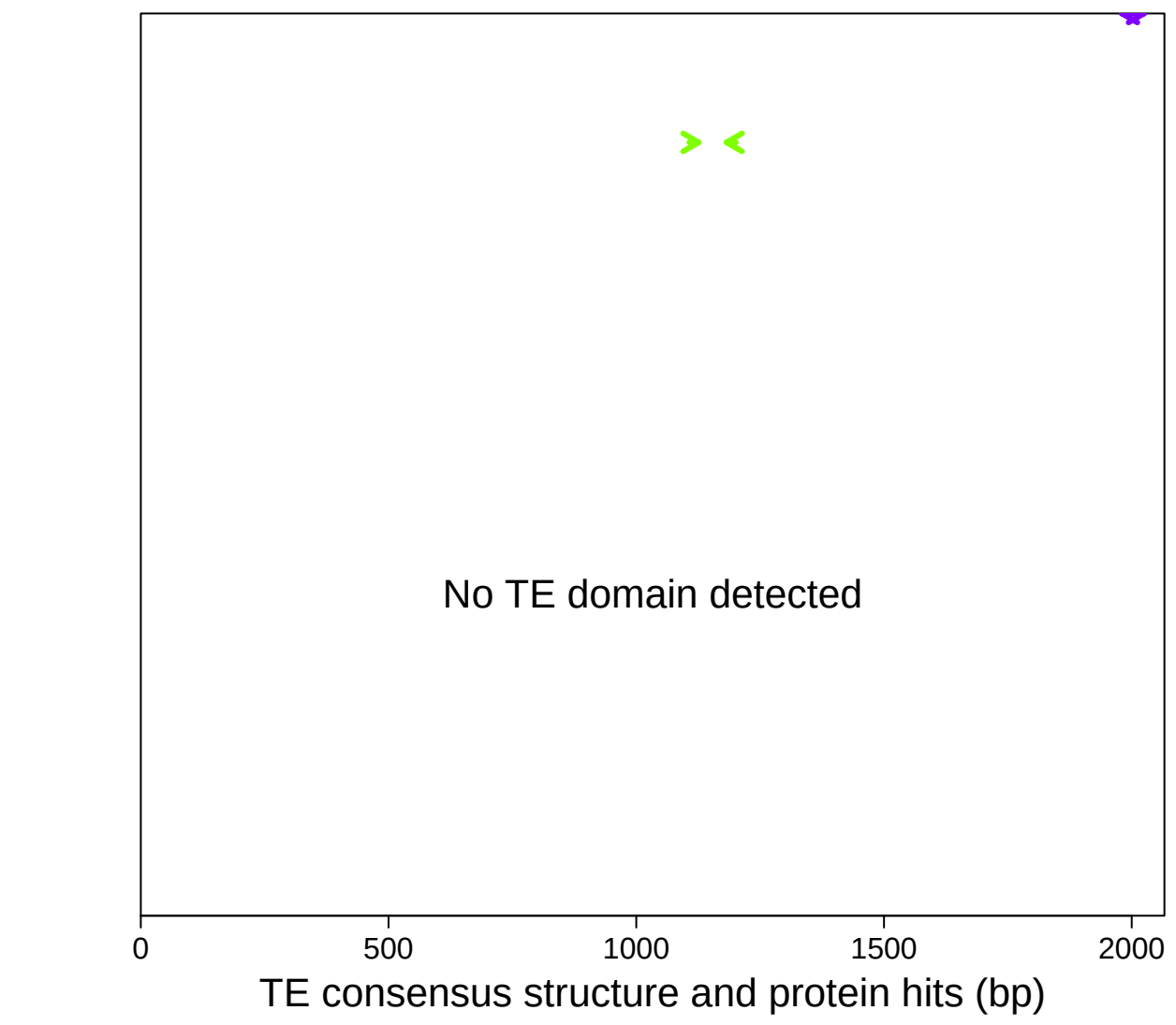
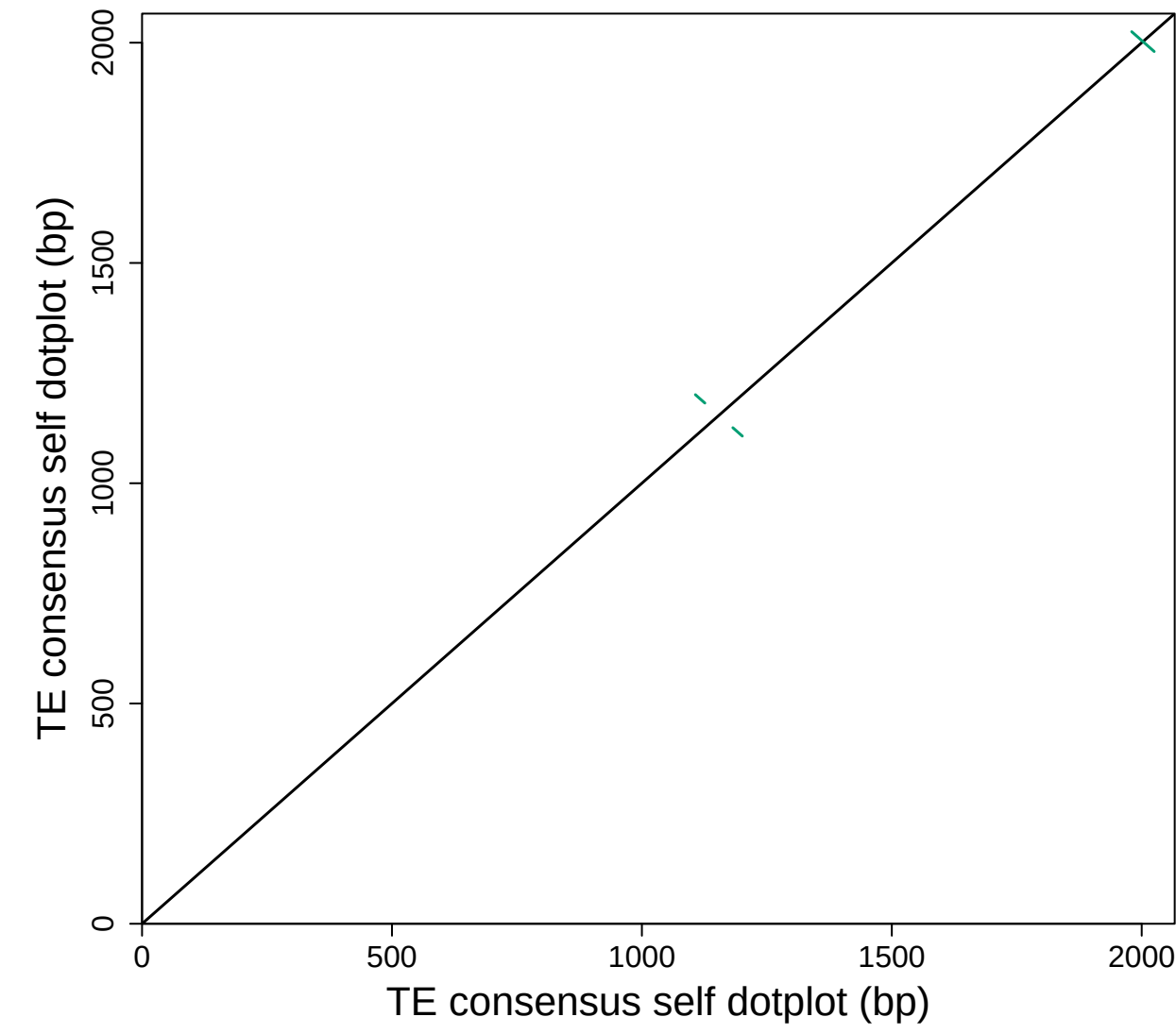
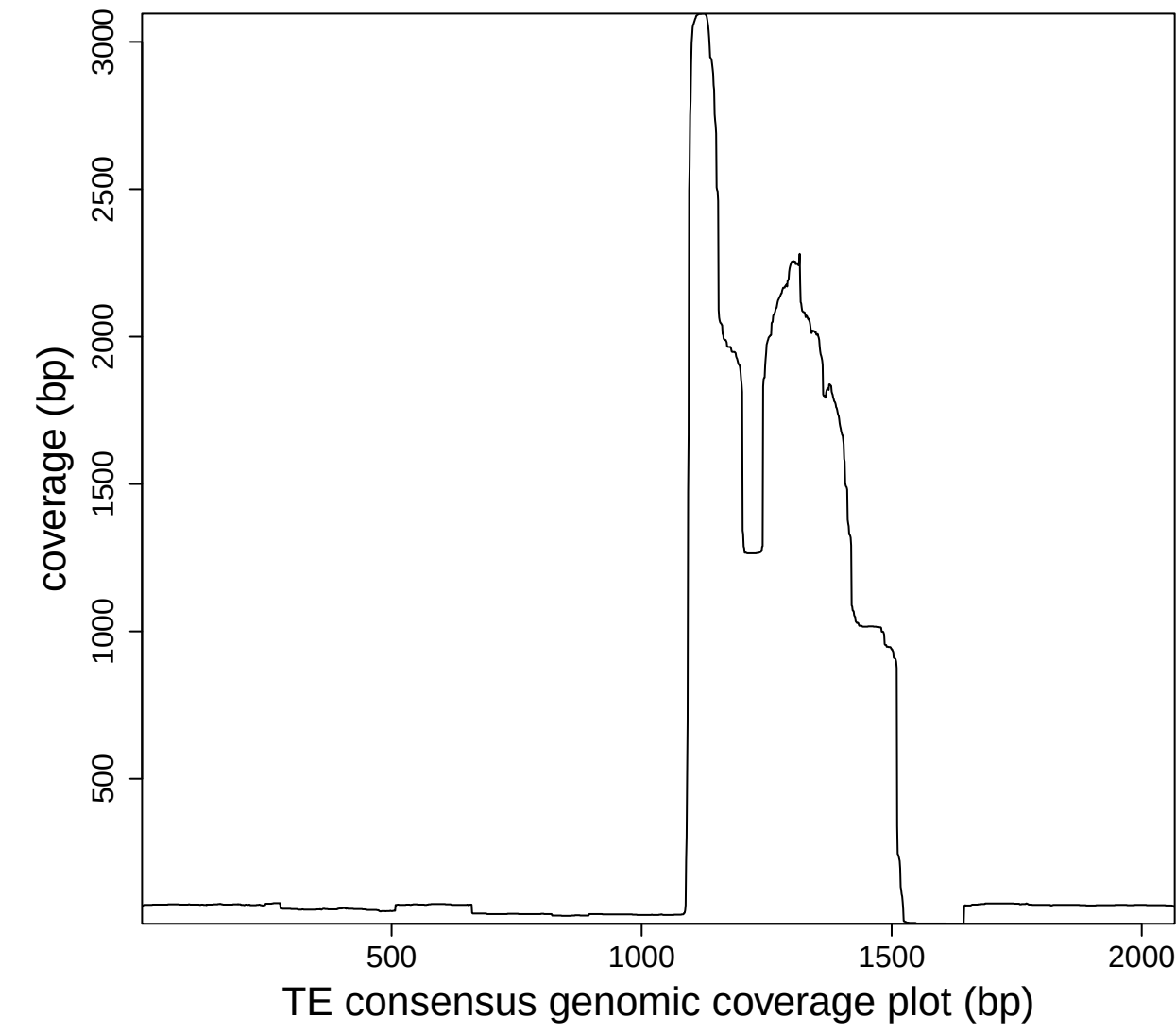
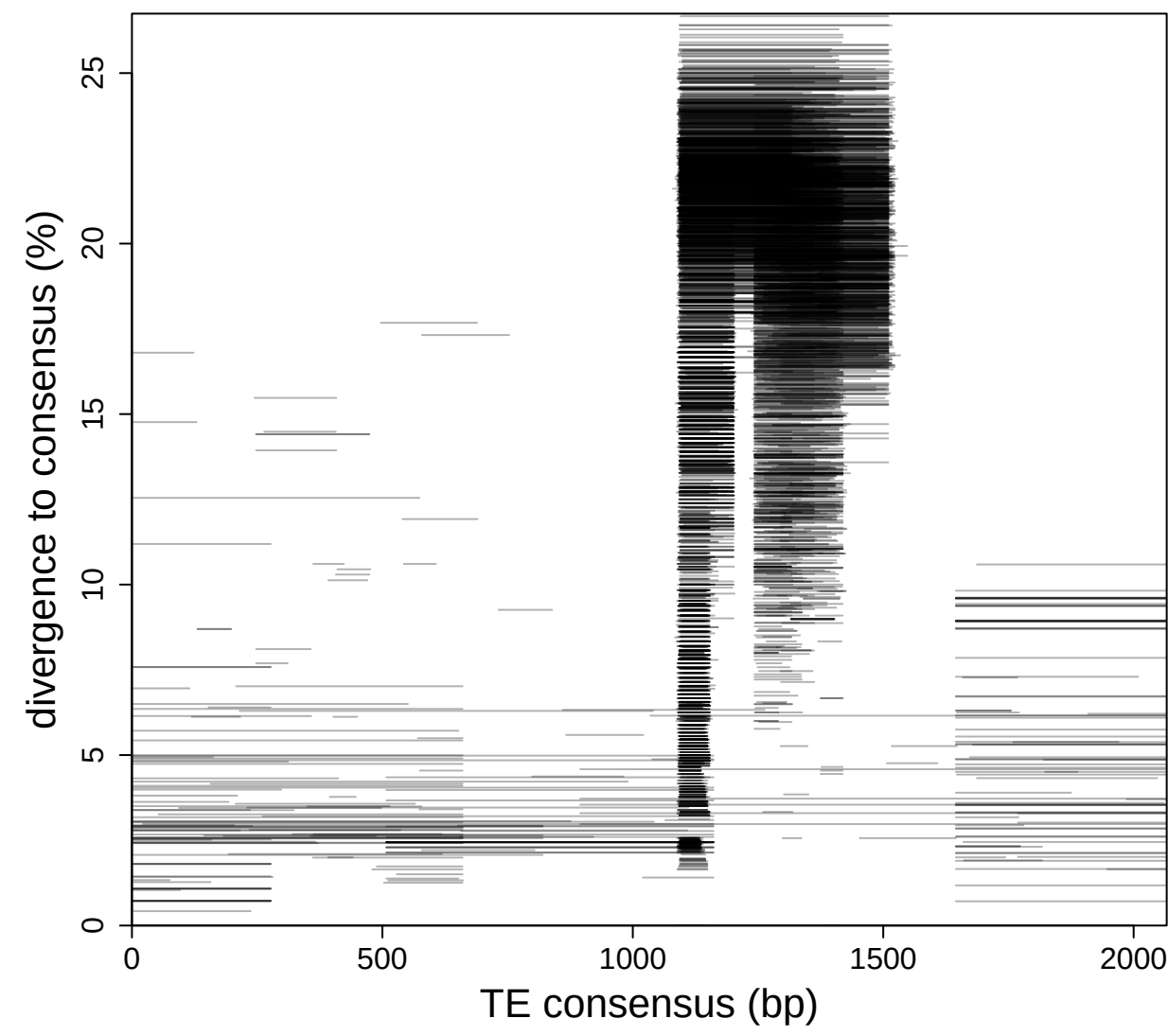
Start crop Point

End crop Point



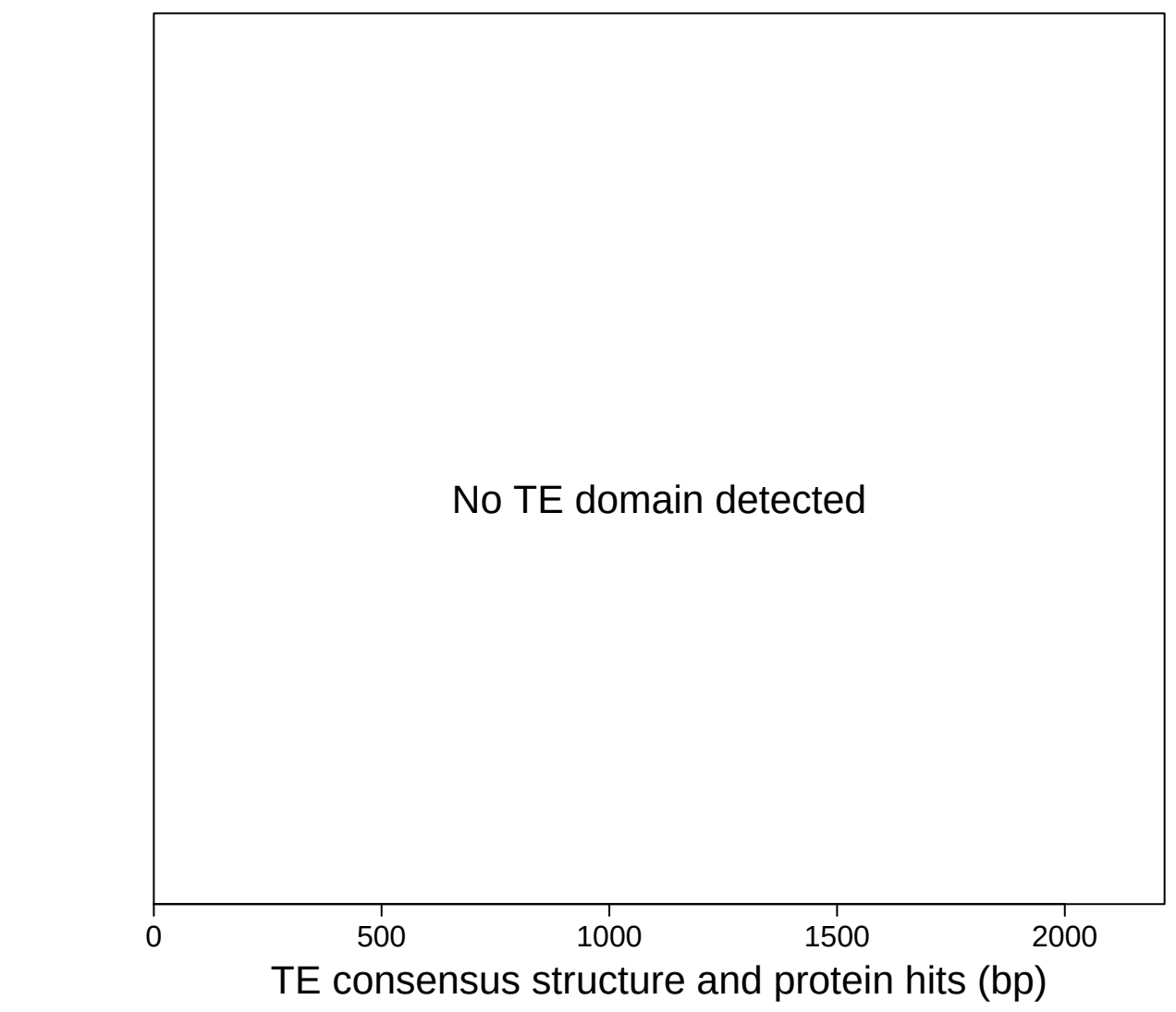
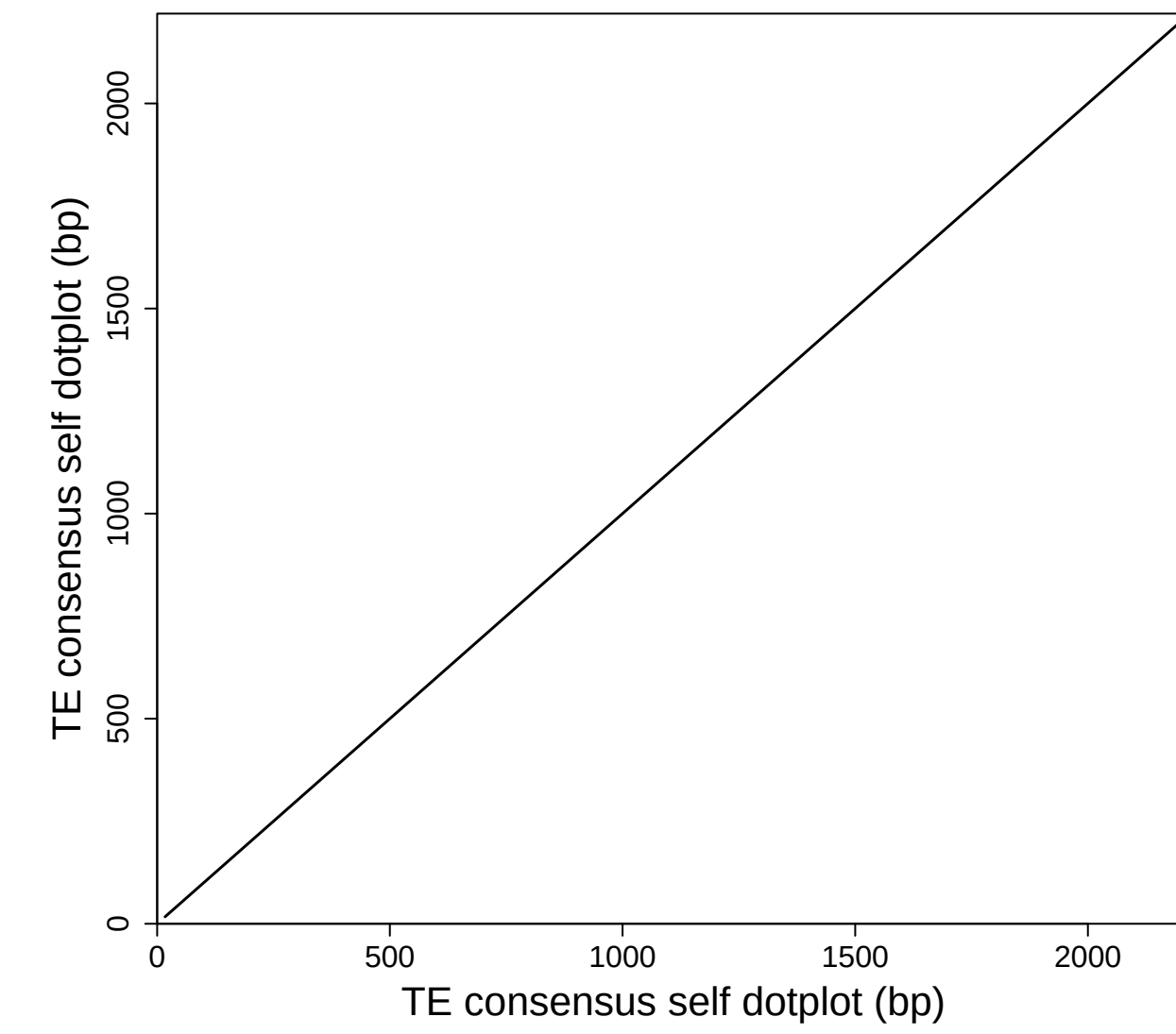
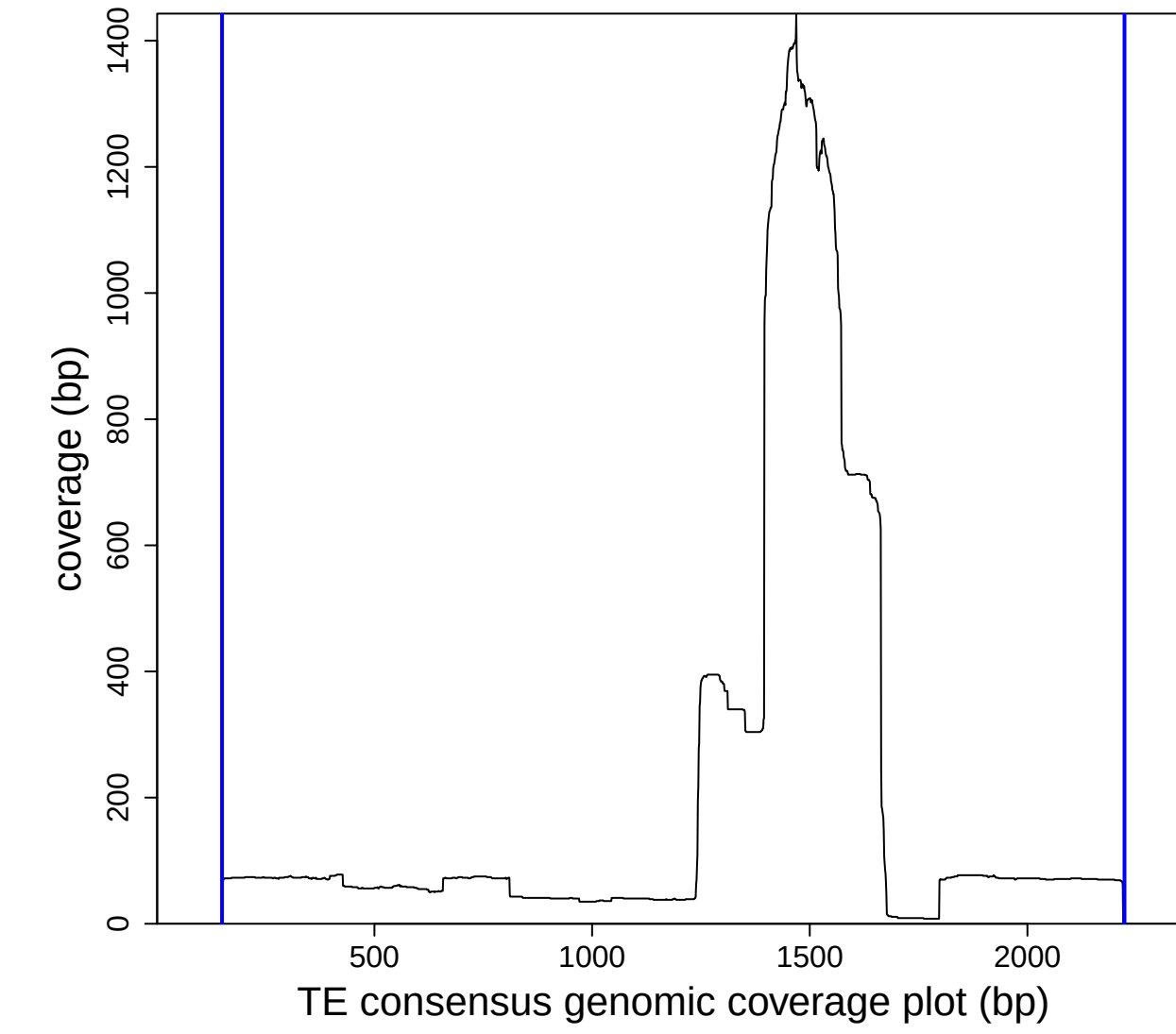
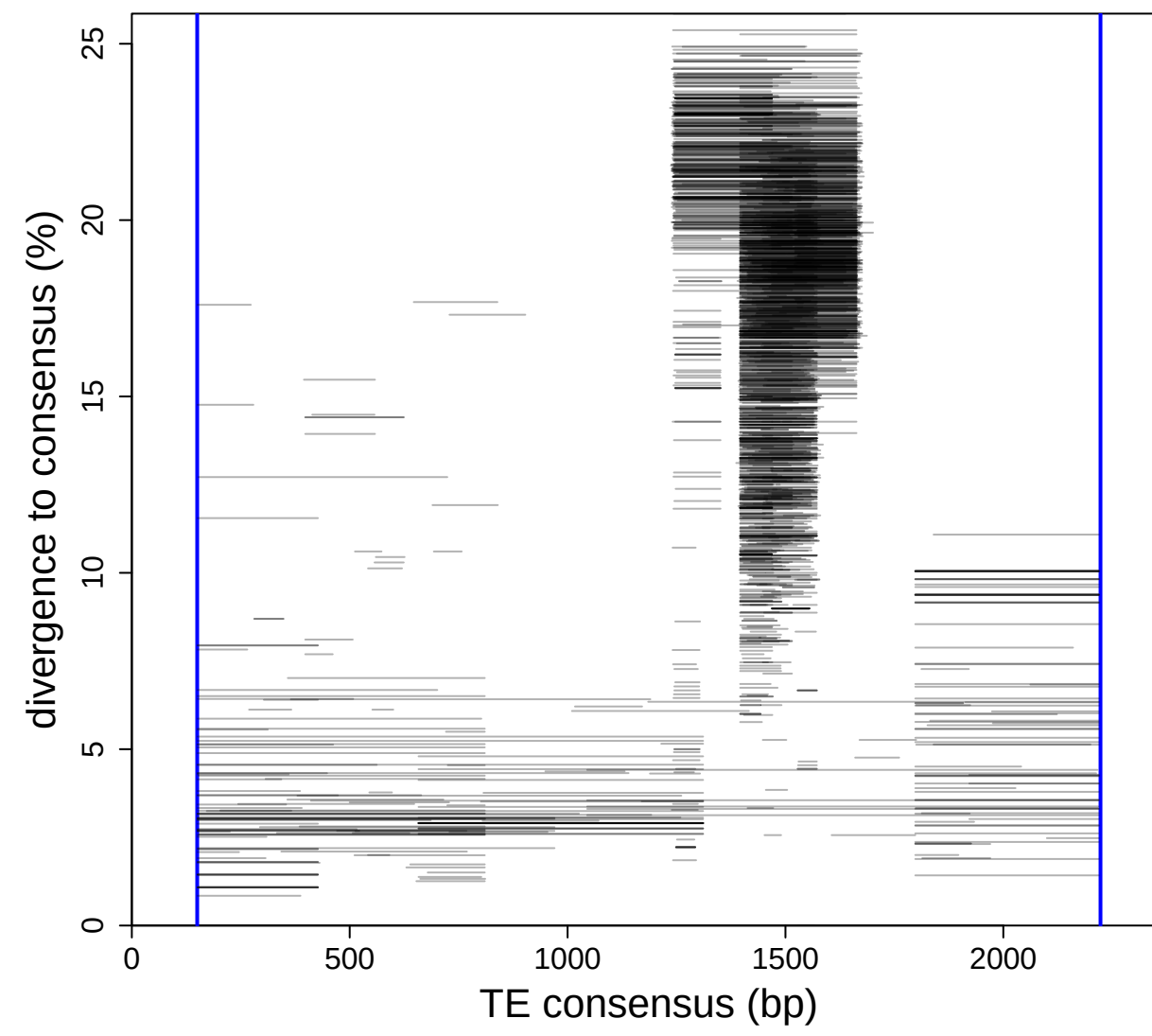
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 2066bp; fragments: 4512; full length: 0 (>=1859.4bp)

After TEtrimmer 2066 bp



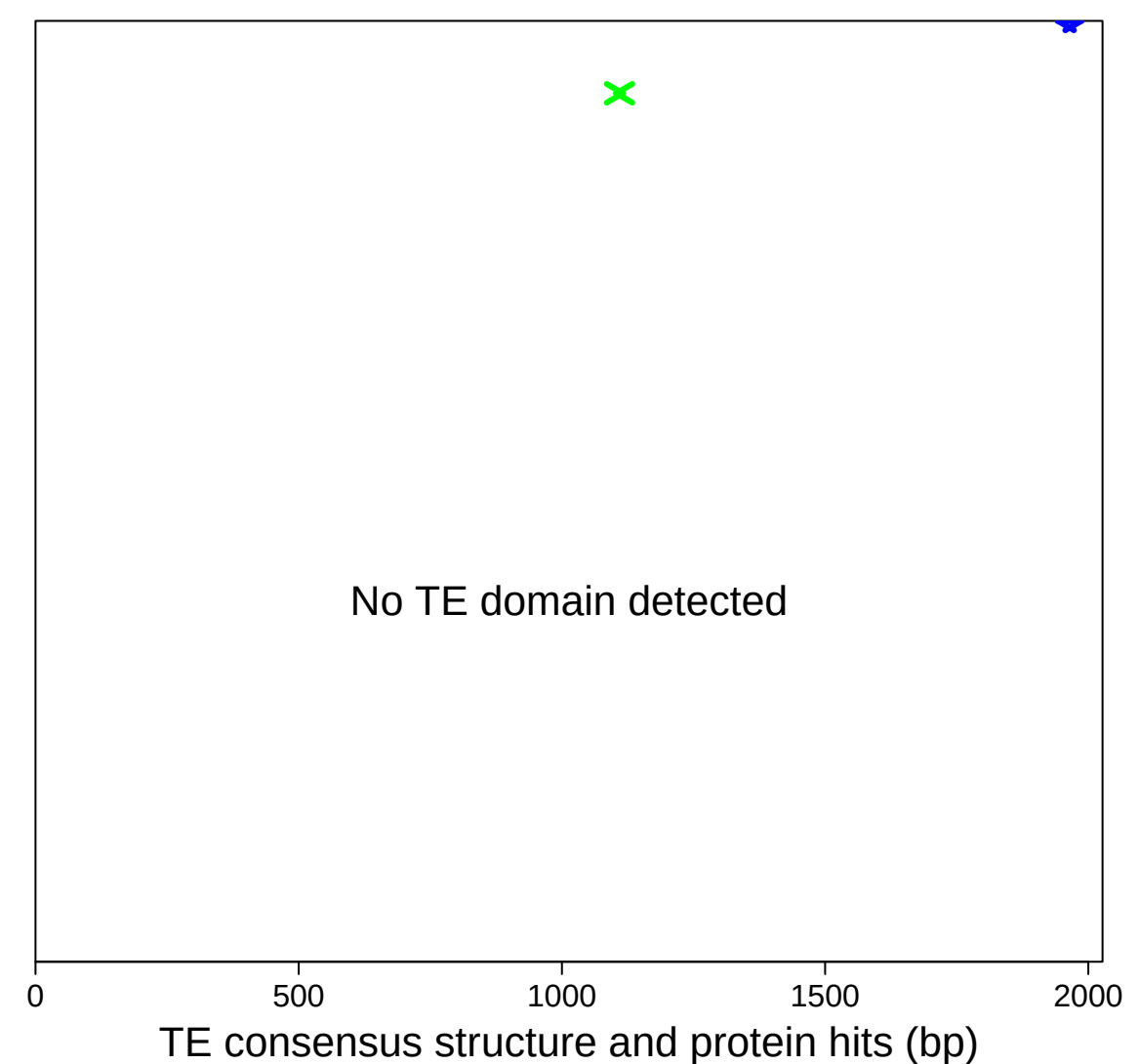
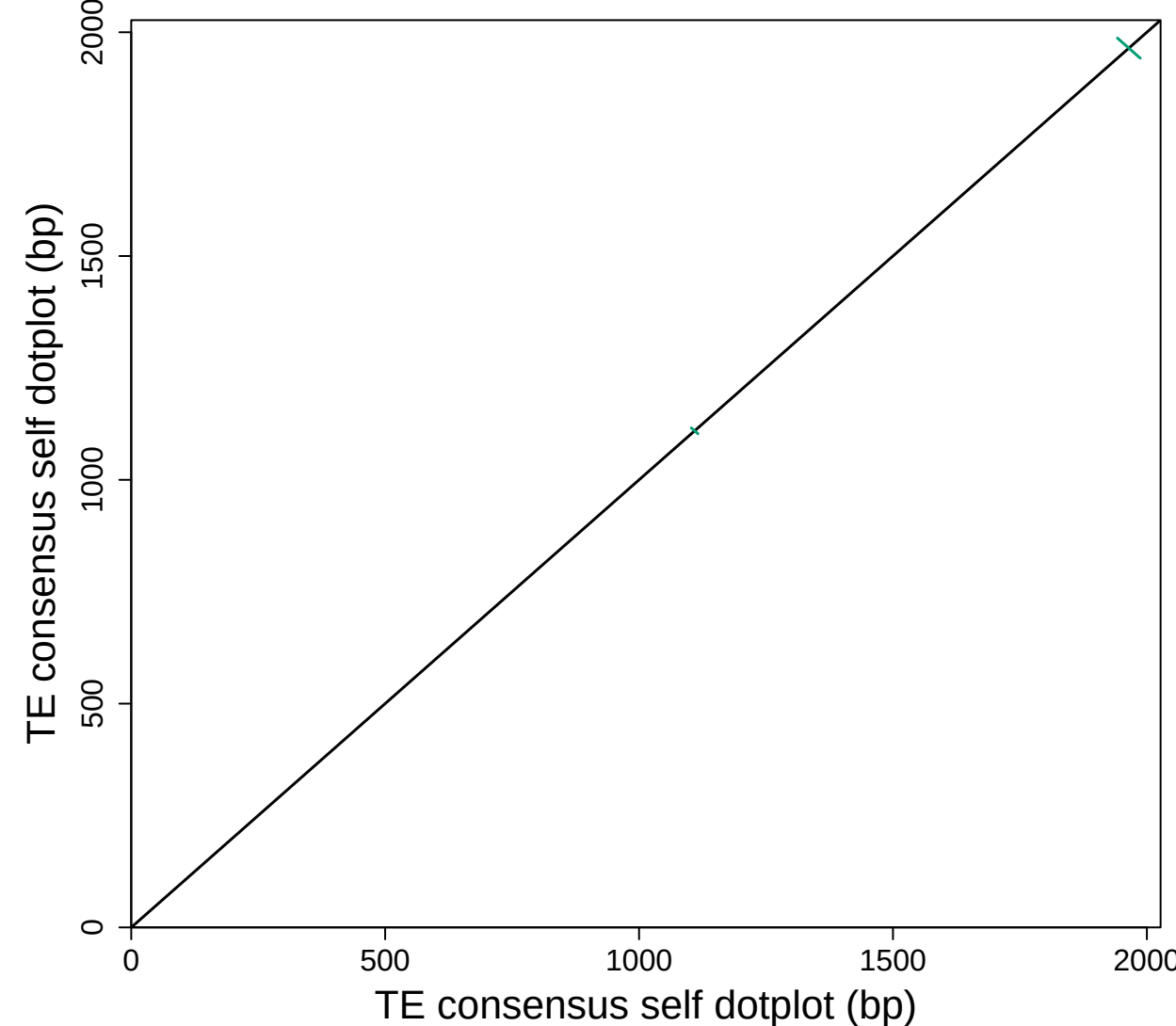
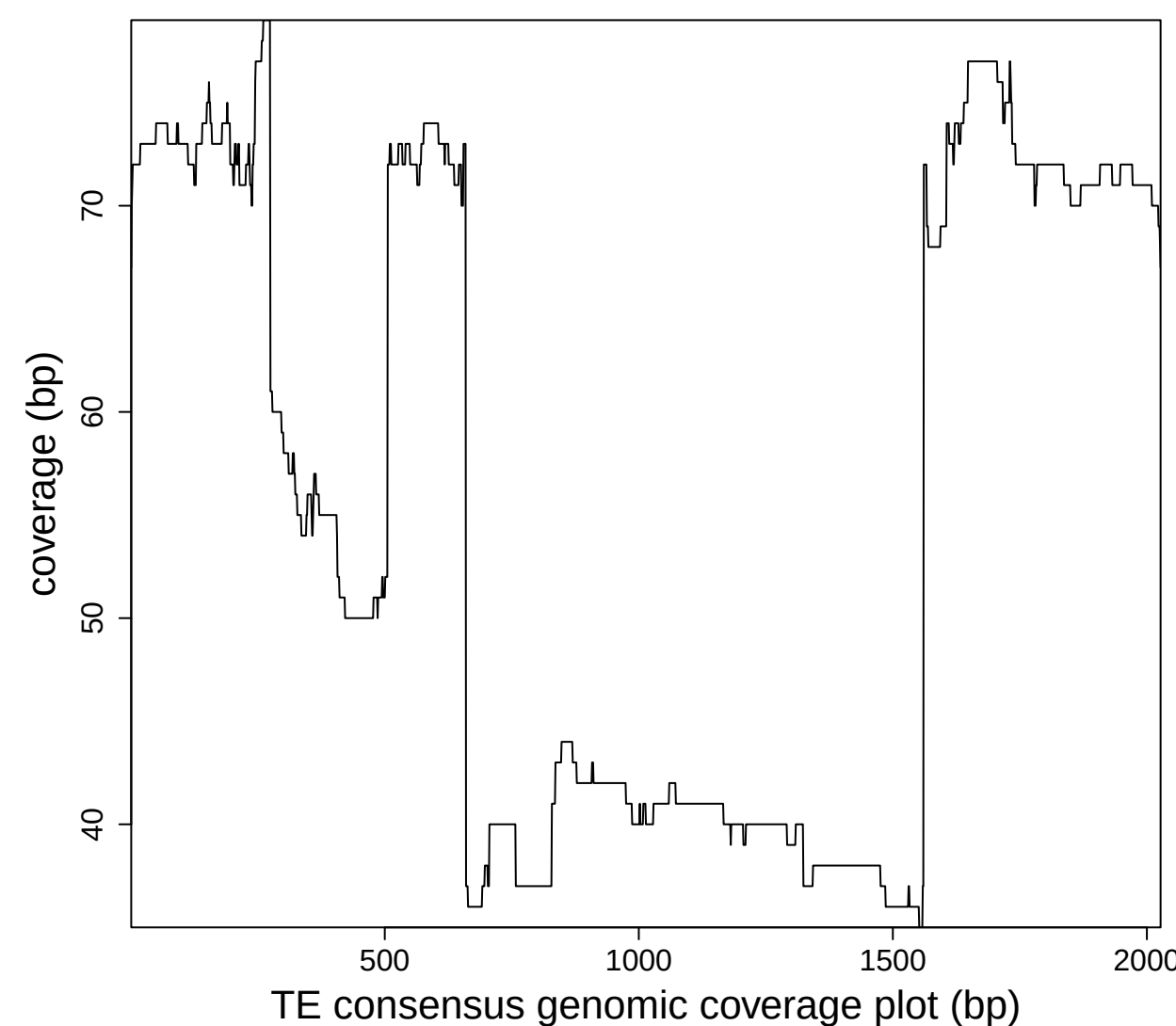
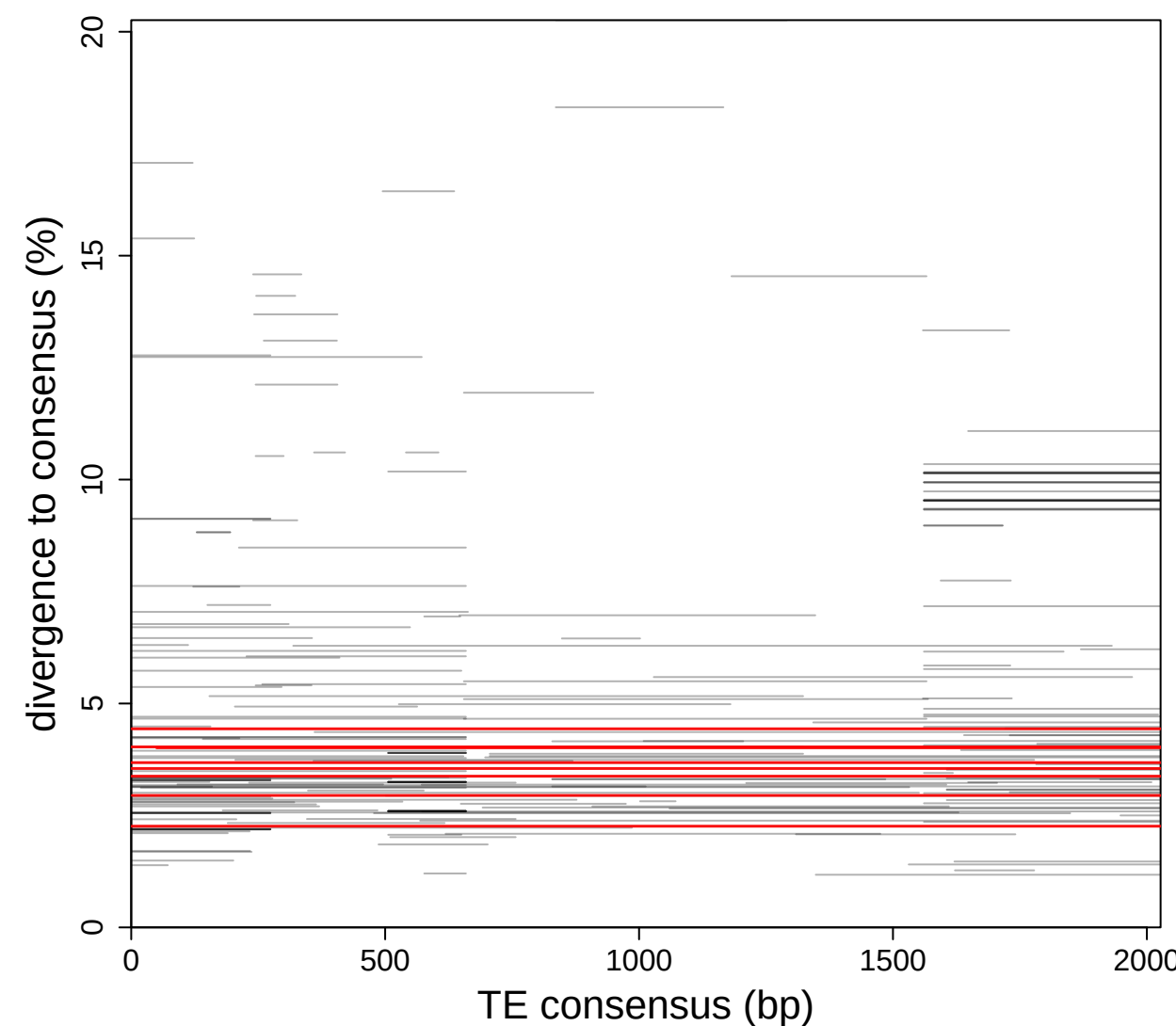
0.fasta.b.bed uf.bed g_3.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 2373bp; fragments: 1892; full length: 0 (>=2135.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

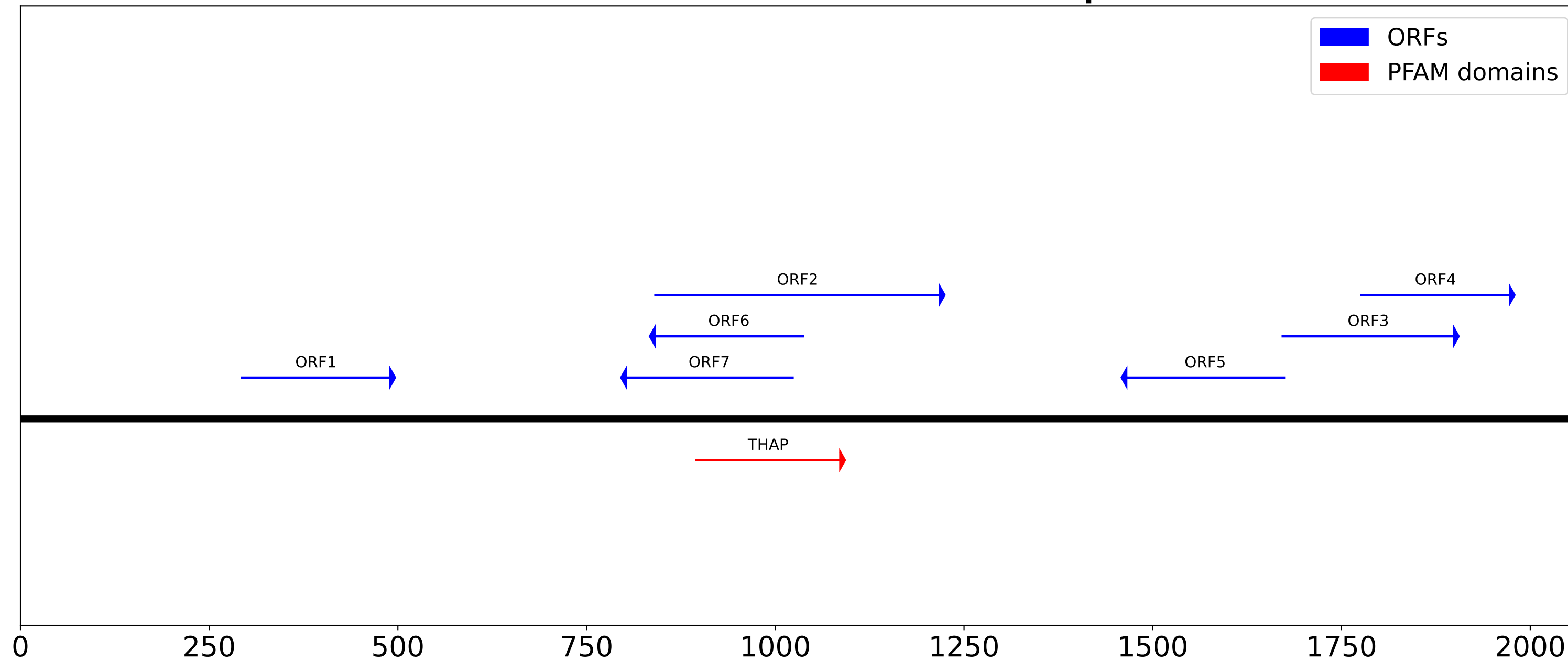


TE: rnd_5_family_1050
size: 2027bp; fragments: 224; full length: 8 (≥ 1824.3 bp)

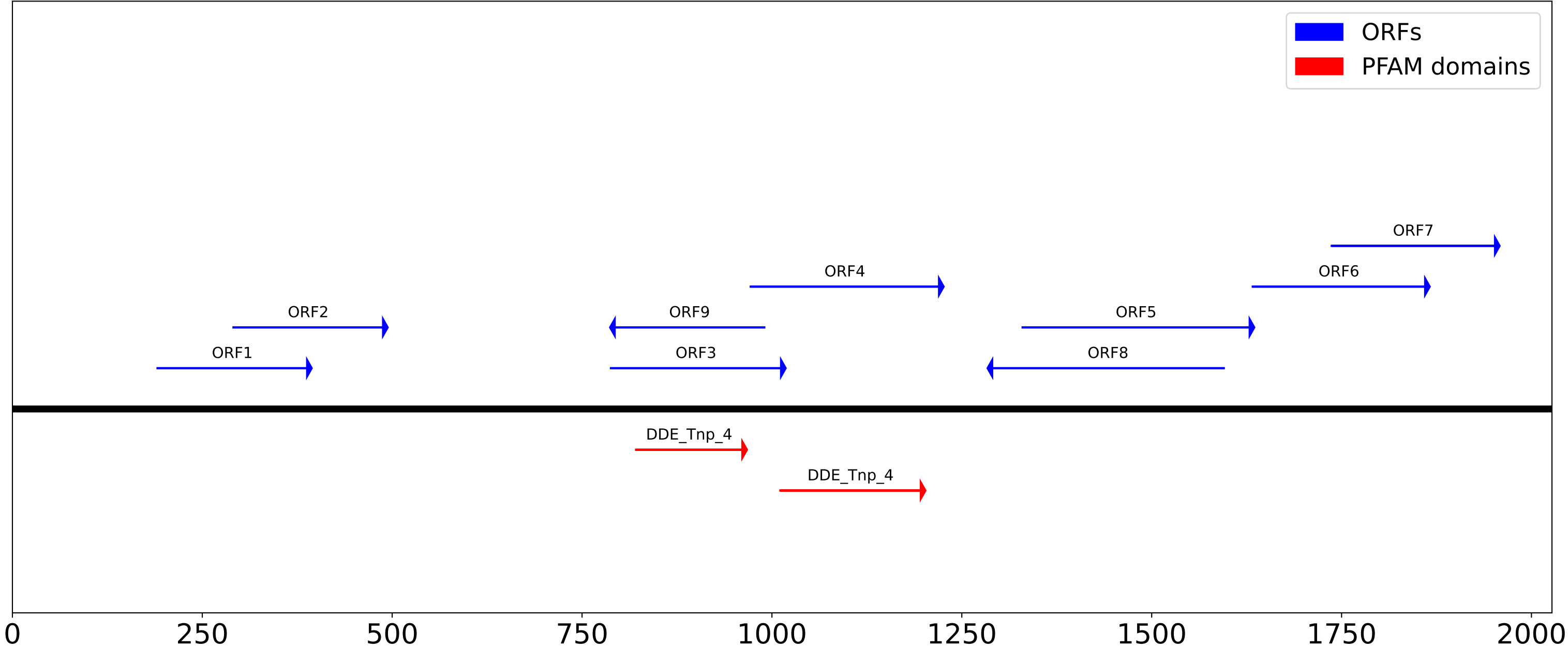
Before TEtrimmer 2027 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 10/10/25)

